Express Mailing Label No. EV139505841US Date of Deposit: February 27, 2003

Attorney Docket No: 21127-501RCE

# IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

APPLICANT: Perrin et al.

09/314,698 SERIAL NUMBER:

EXAMINER: Juliet C. Einsmann

FILING DATE: May 19, 1999

ART UNIT: 1655

FOR: MICRO-ARRAY BASED SUBTRACTIVE HYBRIDIZATION

February 27, 2003

Boston, Massachusetts

MAR 0 6 2003

TECH CENTER 1600/2900

Commissioner for Patents Washington, D.C. 20231

# TRANSMITTAL LETTER

Transmitted herewith for filing in the present application are the following documents:

Amendment and Response;

Petition for Extension of Time (1 pg., in duplicate);

Check #15679 for \$930.00; and

Return Postcard.

If the enclosed papers are considered incomplete, the Mail Room is respectfully requested to contact the undersigned collect at (617) 542-6000, Boston, Massachusetts. The Commissioner is authorized to credit any overpayment or charge any deficiencies to Deposit Account No. 50-0311, Reference No. 21127-501RCE. A duplicate copy of this Transmittal Letter is enclosed.

Respectfully submitted,

Ingrid A. Beattie, Reg. No. 42,306

Attorney for Applicants

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30623

PATENT TRADEMARK OFFICE

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EXAMINER:

Juliet C. Einsmann

FILING DATE:

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ART UNIT:

1634

For:

Micro-Array Based Subtractive Hybridization

RECEIVED

Assistant Commissioner for Patents

Washington, D.C. 20231

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TECH CENTER 1600/2900

#### RESPONSE AND AMENDMENT

In response to the Office Action mailed on August 27, 2002, please amend the application as follows.

# In the Figures:

Please replace the pending figures with the enclosed replacement Figures 1-4 (4 sheets).

### In the Claims:

Cancel claims 27-57.

Add new claims 58-62.

- 58. A method of enhancing the rate of novel gene discovery in a population of previously-uncharacterized nucleic acid molecules, comprising
- a. contacting an first ordered array of nucleic acid samples from said population with an initial complex subtraction probe pool, said probe pool comprising a labeled probe corresponding to a redundant or known sequence;
- b. identifying a nucleic acid, said nucleic acid being characterized by weak or no hybridization to said initial probe pool;